TABLE 1

Genetic Epidemiology and Core Public Health Functions in the

Continuum from Genes to Public Health

| Step | Description of activities | Disease/gene examples |
|----------------------------------------|-------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------|
| GENETIC TECHNOLOGY HUMAN GENOME PROJEC | Gene mapping and linkage T studies in high risk families | 50,000-100,000 genes such as BRCA1 in breast cancer |
| V ASSESSMENT | GENETIC EPIDEMIOLOGY | ApoE E4 allele and Alzheimer's disease |
| V POLICY DEVELOPMENT | When and how genetic tests are to be applied in public health programs | Screening for various genes |
| V ASSURANCE | Development of public health genetic programs, evaluation of prevention effectiveness, quality assurance | |

TABLE 2

Genetic Epidemiology and Assessment of the Role of Genetic Factors in Disease

| Ası | pects | Examples | | |
|-----------|-------------------------------------------------------------|------------------------------------------------------------------------------------|--|--|
| <u>I.</u> | Population studies | | | |
| Α. | Prevalence of susceptibility alleles in various populations | Studies of the frequency of BRCA1 mutations in different ethnic groups | | |
| В. | Determinants of mutations in various populations | Studies of risk factors for chromosomal anomalies such as Down syndrome | | |
| C. | Association between genetic traits and diseases | Studies of ApoE-E4 allele in Alzheimer's disease in the population | | |
| <u>II</u> | . Family studies | | | |
| Α. | Familial aggregation of diseases | Recurrence risks of birth defects after an an affected pregnancy | | |
| В. | Causes of familial aggregation of disease | Studies of genetic and environmental factors in the recurrence of various diseases | | |
| C. | Establishing genetic modes of inheritance | Segregation and linkage analysis in families | | |

TABLE 3

Gene-Environment Interaction Analysis in a Case-Control Study

| Exposure | Suscep- tibility genotype | Cases | Controls | Odds Ratio |
|-----------|---------------------------------|-----------------|-----------------|---------------------------------------|
| (1=presen | t, 0=absent) | | | |
| 0 | 0 | A ₀₀ | B ₀₀ | $OR_{00} = 1.0$ |
| 0 | 1 | A ₀₁ | B ₀₁ | $OR_{01} = A_{01}B_{00}/A_{00}B_{01}$ |
| 1 | 0 | A_{10} | B ₁₀ | $OR_{10} = A_{10}B_{00}/A_{00}B_{10}$ |
| 1 | 1 | A_{11} | B ₁₁ | $OR_{11} = A_{11}B_{00}/A_{00}B_{11}$ |

Case-only odds ratio $OR_{ca}=A_{11}A_{00}/A_{10}A_{01}=(OR_{11}/OR_{10}OR_{01})OR_{co}$ Where $OR_{co}=B_{11}B_{00}/B_{10}B_{01}$ (control-only odds ratio)

TABLE 4

Case-Control Analysis of the Interaction Between Maternal
Cigarette Smoking, Transforming Growth Factor Alpha
Polymorphism, and the risk for cleft palate. Adapted from Hwang
et al. (11)

| Smoking P | TaqI olymorphism | Cases | Controls | Odds Ratio | 95% C.I. |
|--------------|---------------------|-------|----------|---------------|----------|
| _ | _ | 36 | 167 | 1.0 | Referent |
| - | + | 7 | 34 | 1.0 | 0.3-2.4 |
| + | - | 13 | 69 | 0.9 | 0.4-1.8 |
| + | + | 13 | 11 | 5.5 | 2.1-14.6 |
| | | | | | |

Crude odds ratios are presented. Odds ratio based on a case-only study is 5.1 (95% C.I. 1.5-18.5) (13 * 36)/(13 * 7)

TABLE 5
Characteristics of Nontraditional Case-Only Studies

| Feature (| Case-Only | Case-Parental Control | Affected Relative-pair |
|-------------------------|-------------------------------------------------------------------------------|--------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------|
| Study Subjects | Cases | Cases and their parents | Second case in family, proband, and parents |
| 'Controls' | None | Expected genotype distribution based on parental genotypes | Expected distribution of alleles with Mendelian transmission |
| Assessment | Departure from multiplicative relation between exposure and genotype | Association between genotype and disease Also departure from multipli- | Linkage between locus and disease Also departure from multipli- |
| | | cative relation | cative relation |
| Assumptions | s Independence between genotype and exposure | Mendelian tranmission | Mendelian transmission |
| Strengths & limitations | Simple. Cannot assess effects of exposure or genotype. Linkage disequilibrium | Requires one or both parents. Cannot assess exposure effects. Linkage disequilibrium | Need families with 2 or more cases. Cannot assess exposure Cannot assess specific |

TABLE 6

Gene-Environment Interaction Analysis in the Context of a CaseParental Control Study: Analysis of Nontransmitted Alleles

Susceptibility genotype

Exposure: Absent

Cases

| | | Present | Absent |
|----------------------------------|--------|---------|--------------------------------|
| Parental non-transmitted all | - | T_0 | U_0 |
| | Absent | V_0 | W_0 |
| Odds Ratio (amo unexposed) | | 1 | U ₀ /V ₀ |

Exposure: Present

Cases

| | | | Present | Absent | |
|------------|---------|--|----------------|----------------|--|
| Parental | Present | | \mathtt{T}_1 | \mathtt{U}_1 | |
| non-trans- | | | | | |

mitted alleles

| | Absent | V_1 | W_1 |
|------------|--------|-------|-----------|
| | | | |
| | | | |
| Odds | | 1 | U_1/V_1 |
| Ratio (amo | ong | | |
| exposed) | | | |

TABLE 7

Gene-Environment Interaction Analysis in the Context of an Affected Sib-Pair Study

| Alleles with | Unexposed case | Exposed case | Expected | Odds Ratio (unexposed) | Odds Ratio (exposed) |
|-----------------|-------------------|-----------------|----------|------------------------------|----------------------------|
| 0 | A ₀₀ | A ₀₁ | 0.25 | 1.0 | 1.0 |
| 1 | A ₁₀ | A ₁₁ | 0.50 | $A_{10}/2A_{00}$ | $A_{11}/2A_{01}$ |
| 2 | A_{20} | A_{21} | 0.25 | A_{20}/A_{00} | A_{21}/A_{01} |
| | | | | | |

TABLE 8

Incorporating a Familial Analysis of Reconstructed Cohorts into a Case-Control Study

| Disease in a Relative | Disease in Index Persons | | |
|--------------------------|-----------------------------|---------|--|
| | Case | Control | |
| Yes | A_1 | A_0 | |
| No | B_1 | B_0 | |
| Total | N_1 | N_0 | |
| | | | |

Total of case relatives is N_1 and control relatives $N_0.$ Disease proportion in case relatives A_1/N_1 Disease proportion in control relatives A_0/N_0 Risk ratio $(A_1/N_1)/(A_0/N_0)$

TABLE 9
Linkage analysis in an Epidemiologic Study Design

| Alleles ibd w probands | Cohort study | | Case-control study | | |
|------------------------------|-----------------|---------------|--------------------|----------|--------------------------|
| probanas | Disease Risk | Risk ratio | Recurrent cases | Controls | Odds ratio |
| | | | | | |
| 0 | R_0 | 1.0 | A_0 | B_0 | 1.0 |
| 1 | R_1 | RR_1 | A_1 | B_1 | $\mathrm{A_1B_0/A_0B_1}$ |
| 2 | R_2 | RR_2 | A_2 | B_2 | A_2B_0/A_0B_2 |
| | | | | | |